

**IN THE CLAIMS:**

Please amend the claims as follows:

- B7  
C10
13. (Amended) A peptide of the sequence:  
 $X^1$ m-Leu-Arg-Ile-Val-Gln-Cys-Arg-Ser-Val-Glu-Gly-Ser-Cys-Gly-Phe- $X^2$ n  
 (SEQ ID NO: 2)  
 wherein  $X^1$  and  $X^2$  are each selected from the group consisting of L- or D- Arg, His, Lys and Tyr, and m and n are each 0, 1, 2 or 3 with the proviso that at least m or n is 1;  
 a cyclic disulfide thereof or an organic or inorganic acid addition salt thereof.
14. (Amended) A peptide of the sequence:  
 $Y^1$ -Leu-Arg-Ile-Val-Gln-Cys-Arg-Ser-Val-Glu-Gly-Ser-Cys-Gly-Phe (SEQ ID NO: 3)  
 wherein  $Y^1$  is selected from the group consisting of the desamino form (H), acetyl ( $CH_3CO-$ ) and other acyl groups;  
 a cyclic disulfide thereof or an organic or inorganic acid addition salt thereof.
15. (Amended) A peptide of the sequence: (SEQ ID NO: 4)  
 Leu-Arg-Ile-Val-Gln-Cys-Arg-Ser-Val-Glu-Gly-Ser-Cys-Gly-Phe- $Y^2$   
 wherein  $Y^2$  is selected from the group of  $CONH_2$  and alkyl amide groups;  
 a cyclic disulfide thereof or an organic or inorganic acid addition salt thereof.
16. (Amended) A peptide which is selected from the group consisting of:

Ref No.  
 (SEQ ID NO: 15)  
 9502

**STRUCTURE**

Leu Arg Ile Val Gln Pen Arg Ser Val Glu Gly Ser Pen Gly

Phe

(SEQ ID NO: 8)

9405

$CH_3CO-$  Leu Arg Ile Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly

Phe

B7

(SEQ ID NO: 12)  
 9410 H - Leu Arg Ile Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly  
 Phe

(SEQ ID NO: 7)  
 9404 Leu Arg Ile Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly  
 Phe - CONH<sub>2</sub>

(SEQ ID NO: 10)  
 9407 Leu Arg Ile Val Gln Cys Lys Ser Val Glu Gly Ser Cys Gly  
 Phe

(SEQ ID NO: 11)  
 9408 Leu Arg Ile Val Gln Cys Lys Ser Val Glu Gly Ser Cys Gly  
 Phe

|\_\_\_\_\_|(amide bond)

9604 Tyr Leu Arg Ile Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly  
 Phe

(SEQ ID NO: 19)  
 9605 Lys Leu Arg Ile Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly  
 Phe

(SEQ ID NO: 20)  
 9618 Lys Lys Leu Arg Ile Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly Phe

(SEQ ID NO: 33)  
 9607 Ala Arg Ile Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly Phe

(SEQ ID NO: 22)  
 9606 Leu Lys Ile Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly Phe

(SEQ ID NO: 21)  
 9608 Leu Arg Ala Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly Phe

(SEQ ID NO: 23)  
 9403 Leu Arg Lys Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly Phe

(SEQ ID NO: 6)  
 9609 Leu Arg Ile Ala Gln Cys Arg Ser Val Glu Gly Ser Cys Gly Phe

(SEQ ID NO: 24)  
 9610 Leu Arg Ile Val Ala Cys Arg Ser Val Glu Gly Ser Cys Gly Phe

(SEQ ID NO: 24)  
 9612 Leu Arg Ile Val Gln Cys Arg Ala Val Glu Gly Ser Cys Gly Phe

(SEQ ID NO: 27)  
 9613 Leu Arg Ile Val Gln Cys Arg Ser Ala Glu Gly Ser Cys Gly Phe

See C10 cont